



# A simulation comparison of spatiotemporal and spatially-implicit size-structured models for northern shrimp and snow crab

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# Outline

- Spatially-implicit/ Spatially-stratified/ Spatiotemporal models
- Spatiotemporal modeling framework (snow crab/northern shrimp)
- Simulation experiments
  - recover spatial patterns/unbiased estimates of spatially-aggregated population quantities
  - implicitly accounts for movement processes
  - outperforms spatially-implicit models

# Why spatiotemporal model ?

- Heterogeneous and complex spatial structure – population and fishery
- Spatially-implicit models – biased estimates of population quantities
- Spatially-stratified models
  - spatial strata/movement of individuals among strata

**Correlations - process errors/fishery patterns**

**Spatial correlation (either based on adjacency or distance )**

# Spatiotemporal population model

- Combines theory and methods from population dynamics and geostatistics
- Assume population density varies continuously across space

$$x(s_i) \sim N \left( \frac{1}{|n_i|} \sum_{j \in n_i} x(s_j), \sigma^2 \right)$$

- Joint distribution for density at all locations
- Expand to account for size-structured population dynamics

# Objectives

- Development – estimating population dynamics at a fine spatial scale
- Demonstration – two species (snow crab and northern shrimp)
- Comparison – spatiotemporal model vs. spatially-implicit model
- Evaluation – effect of sample size

# Why size-structured models?

- Advantages:
  - Requires no ability to age animals (shrimps, crabs, lobsters)
  - Uses the data actually available
  - Vulnerability / maturity are often functions of size and not age

# Process model

*Abundance at size (n) for a given location s and time t*

$$\mathbf{n}_{s,t+1} = g(\mathbf{n}_{s,t}) \circ e^{\boldsymbol{\varepsilon}_{s,t}}$$

$$\boldsymbol{\Sigma}_t \sim \text{MVN}(\mathbf{0}, \mathbf{R}_{\text{spatial}} \otimes \boldsymbol{\Theta}_L)$$

- Hadamard product (entrywise product)
- $s$  location
- $t$  year
- ⊗ Kronecker product

$\mathbf{n}_{s,t}$  vector of abundances for each of  $l$  size classes

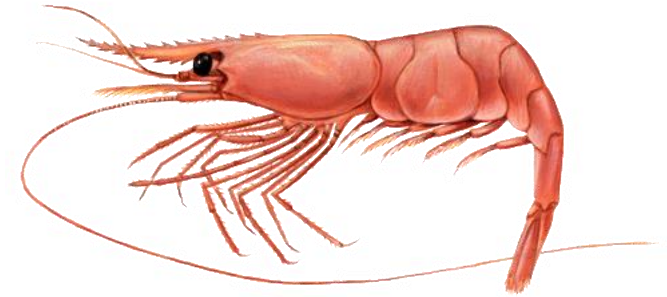
$g()$  function representing population dynamic

$\boldsymbol{\varepsilon}_{s,t}$  vector of random effects (process error)

$\boldsymbol{\Theta}_L$  covariance among size classes ( $l$  by  $l$  matrix  $\mathbf{L}$ )

$\mathbf{R}_{\text{spatial}}$  spatial covariance matrix (covariance between 2 locations follows a Matern function)

# Population dynamic ( $g()$ )



- *Example 1: Gulf of Maine northern shrimp*

$$g(\mathbf{n}_{s,t}) = \mathbf{G}(\mathbf{n}_{s,t-1} \circ \exp(-\mathbf{m}_{s,t-1} - \mathbf{v}f_{s,t-1})) + \mathbf{r}_{s,t}$$

**Initial condition:**

$$g(\mathbf{n}_{s,1}) = \mathbf{r}_{s,t} \circ \exp(\boldsymbol{\varphi})$$

**The predicted harvest per area:**

$$\mathbf{c}_{s,t} = \frac{\mathbf{v}f_{s,t}}{\mathbf{v}f_{s,t} + \mathbf{m}_{s,t}} \circ \left(1 - \exp(-\mathbf{m}_{s,t} - \mathbf{v}f_{s,t})\right) \circ \mathbf{n}_{s,t}$$

$$\log(f_{s,t}) | \log(f_{s,t-1}) \sim \text{N}(\log(f_{s,t-1}), \sigma_f^2)$$



# Population dynamic ( $g()$ )



- *Example 2: Eastern Bering sea snow crab*

$$g(\mathbf{n}_{s,t}^{\text{male}}) = \begin{cases} \mathbf{r}_{s,t} p^{\text{male}} + \mathbf{G}^{\text{male}}(\mathbf{n}_{s,t-1}^{\text{male}} \circ \exp(-\mathbf{m}_{s,t-1} - \mathbf{v}f_{s,t-1}^{\text{male}})) \circ (1 - \mathbf{w}^{\text{male}}), & n = \mathbf{n}^{\lambda} \\ \mathbf{G}^{\text{male}}(\mathbf{n}_{s,t-1}^{\text{male}} \circ \exp(-\mathbf{m}_{s,t-1} - \mathbf{v}f_{s,t-1}^{\text{male}})) \circ \mathbf{w}^{\text{male}} + \mathbf{n}_{s,t-1}^h \circ \exp(-\mathbf{m}_{s,t-1} - \mathbf{v}f_{s,t-1}^{\text{male}}), & n = \mathbf{n}^{\omega} \end{cases}$$

Initial condition:  $g(\mathbf{n}_{s,1}^{\text{male}}) = \mathbf{r}_{s,1} p^{\text{male}} \circ \exp(\boldsymbol{\varphi}_{\text{male}})$

The predicted harvest per area:  $\mathbf{c}_{s,t} = (1 - \exp(-\mathbf{v}f_{s,t}^{\text{male}})) \circ \mathbf{n}_{s,t}^{\text{male}} \circ \exp(-0.5\mathbf{m}_{s,t})$

$$\log(f_{s,t}) | \log(f_{s,t-1}) \sim N(\log(f_{s,t-1}), \sigma_f^2)$$

# Model parameters and estimation

## Fixed effects

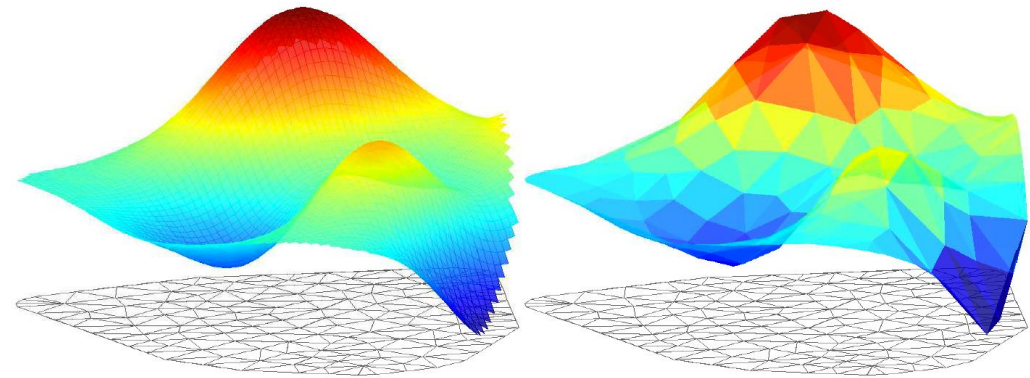
$\Theta_L$	process error covariance (among size classes)
$\kappa$	geostatistical range for correlations
$\mu_t$	average offset of annual recruitment
$\varphi$	initial abundance of each size class
$s$	parameters of selectivity (logistic)
	Parameters of observation model

## Random effects

$r_t^u$	spatial variation in recruitment
$n_t$	spatial variation in density for each size class and year
$f$	fishing mortality of location $s$ over time
	<u>treat density as random, rather than process errors</u>
	$(\varepsilon_t)$

# Gaussian Markov random field (GMRF)

- Continuous spatial process  $\rightarrow$  discretely indexed GMRF
- Matérn covariance function
- Mesh/knot
- SPDE – MVN
- Piecewise constant
- Catch – lognormal
- Survey – lognormal/Poisson-link



Thorson, J.T., Shelton, A.O., Ward, E.J. and Skaug, H.J., 2015. Geostatistical delta-generalized linear mixed models improve precision for estimated abundance indices for West Coast groundfishes. *ICES Journal of Marine Science*, 72(5), pp.1297-1310.

# Input data

## survey data

Size_class	Year	Catch_N	AreaSwept_km2	Vessel	Lat	Lon
1	1	553	3.1	0	60	-174
1	1	629	3.1	0	63.5	-172
1	1	575	3.1	0	58	-170
1	1	618	3.1	0	61.5	-178
1	1	625	3.1	0	64.5	-170
1	1	634	3.1	0	61	-172

- used to create mesh/knots

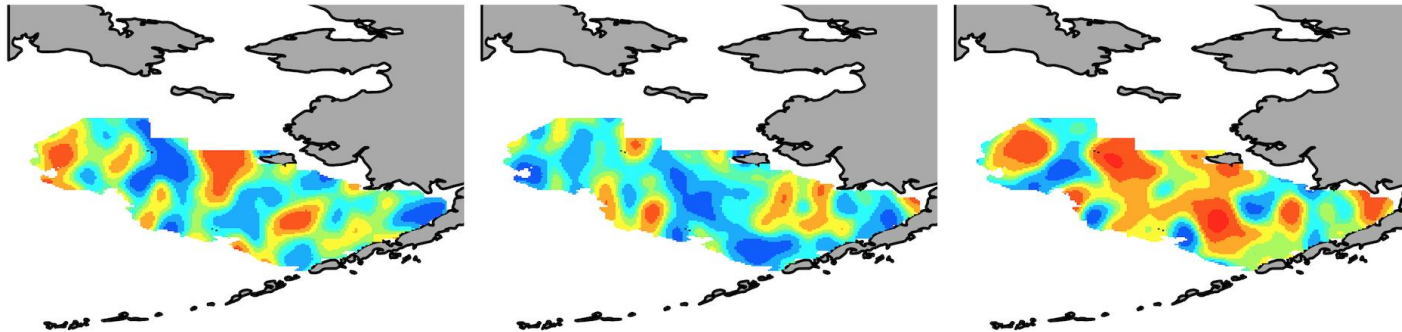
## commercial catch data

X	lat	lon	year	X.1	X.2	X.3	X.4	X.5
1	55.6	-169	1	0.802	2.82	2.32	3.18	7.18
2	56.3	-170	1	0.657	1.83	1.54	2.15	4.94
3	56.3	-170	1	0.662	1.82	1.54	2.16	4.96
4	56.2	-171	1	0.64	1.78	1.5	2.1	4.81
5	56.2	-170	1	0.645	1.8	1.51	2.12	4.85
6	56.2	-170	1	0.646	1.82	1.52	2.13	4.88

- fine scale
- aggregated to knot-level

# Model outputs

- Predicted population density map
- Estimated fishing mortality map
- Predicted catch map
- Estimated covariance of process error



# Simulation – operating model

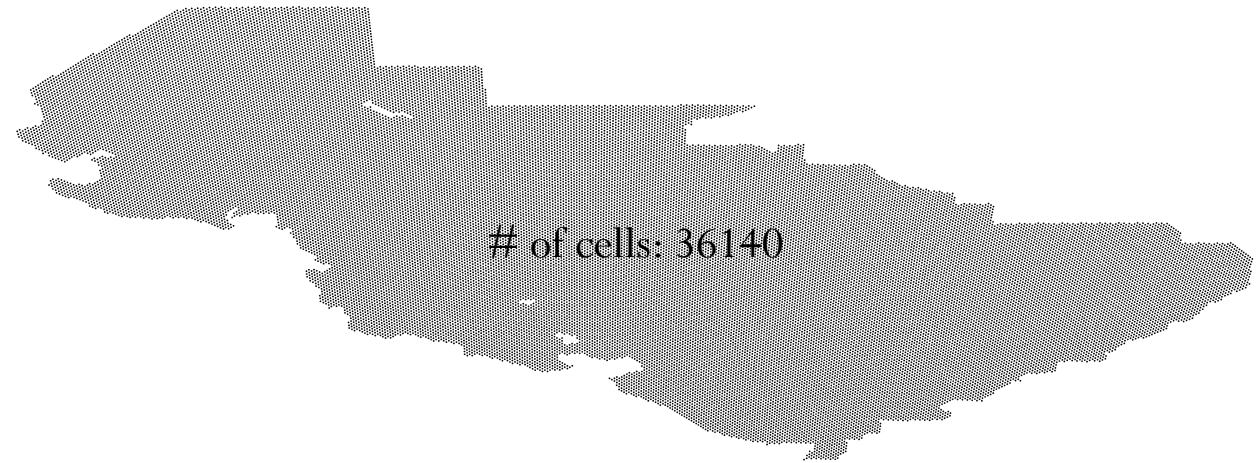
- Dynamics occur at fine scale
- Population dynamics (non-spatial) formulated identically to EM
- Cell-specific parameters (spatially correlated)
- Annual time step
- Movement  $\mathbf{N}_{t+1} = g(\mathbf{M}\mathbf{N}_t) \circ e^{\Sigma_t}$

$$\frac{\partial}{\partial t} \mathbf{n} = \mathbf{N}\mathbf{n}$$

*N* is the matrix of instantaneous movement rates

$$\mathbf{M} \approx \left( \mathbf{I} + \frac{\mathbf{N}\Delta t}{n_{tdiv}} \right)^{n_{tdiv}}$$

*M* is annual movement rates



# Simulation experiments

1. Explore how the spatiotemporal model performs when individual movement processes are modeled explicitly
2. Compare spatially-implicit and spatiotemporal models
3. Evaluate the impact of changing sample size

# Simulation experiments

— Experiment 1: Exploring movement

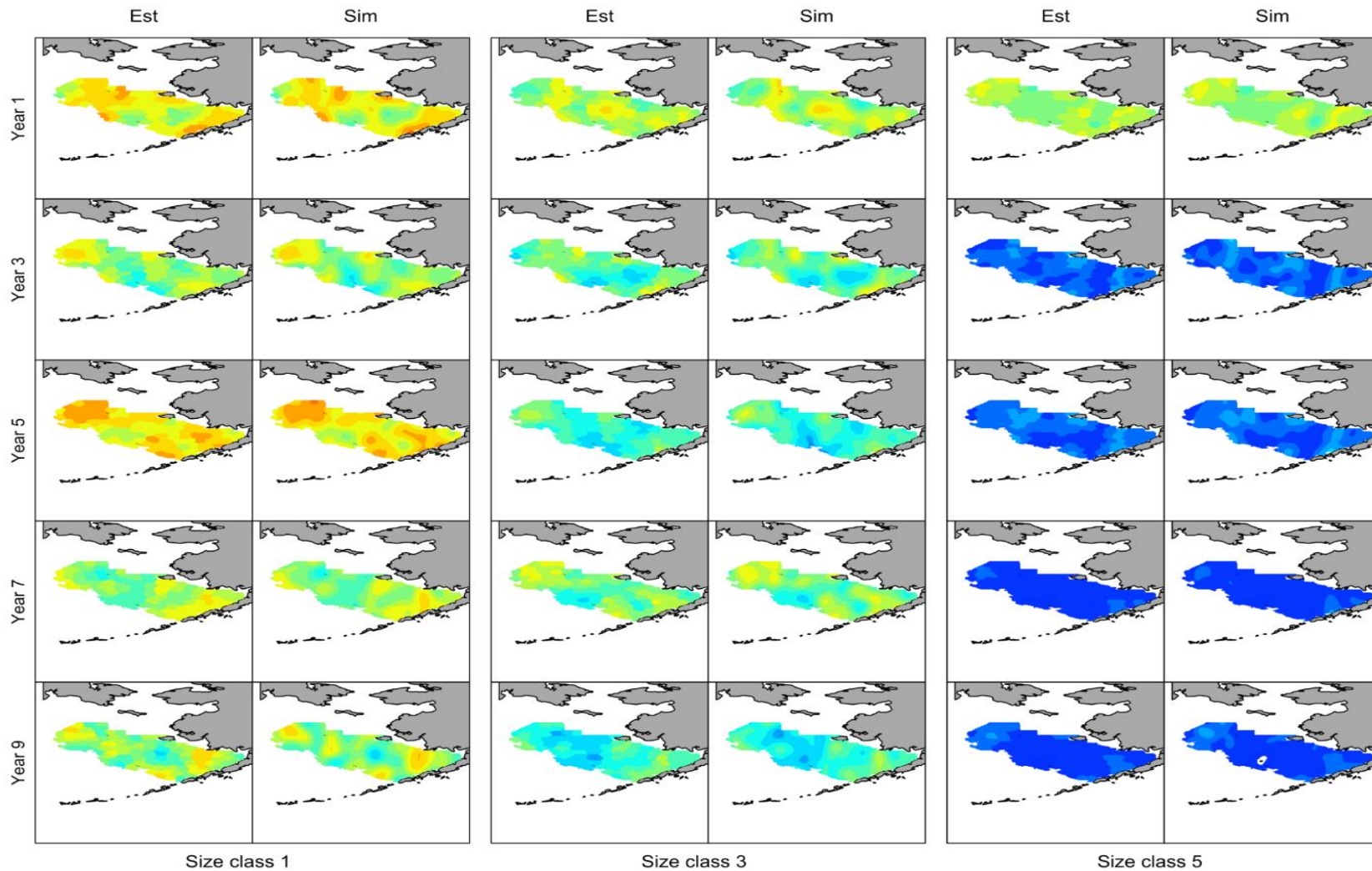


1. No measurement error and no movement in the OM
  2. Same as scenario 1, except there is movement
  3. Both measurement error and movement in the OM
- 200 sites (grid cells) in the OM were randomly sampled each year
  - For each site, total abundance by size class and the total area of the sampled site were recorded
  - Fishery catch-at-size was calculated at each of the 36,140 grids and then aggregated to the knot level as data for the EM
  - For the scenarios with measurement error, we generated 100 replicated data sets with sampling errors, i.e., grid-based survey abundance and fishery catch data were assumed to be lognormally distributed



# Simulation experiments

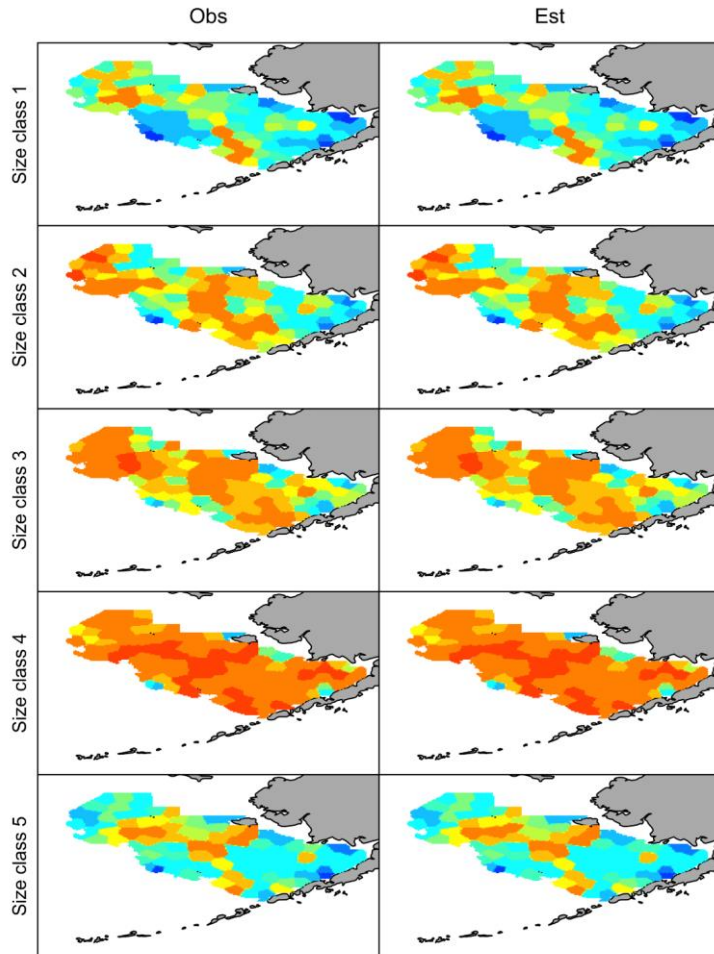
— Experiment 1: Exploring movement



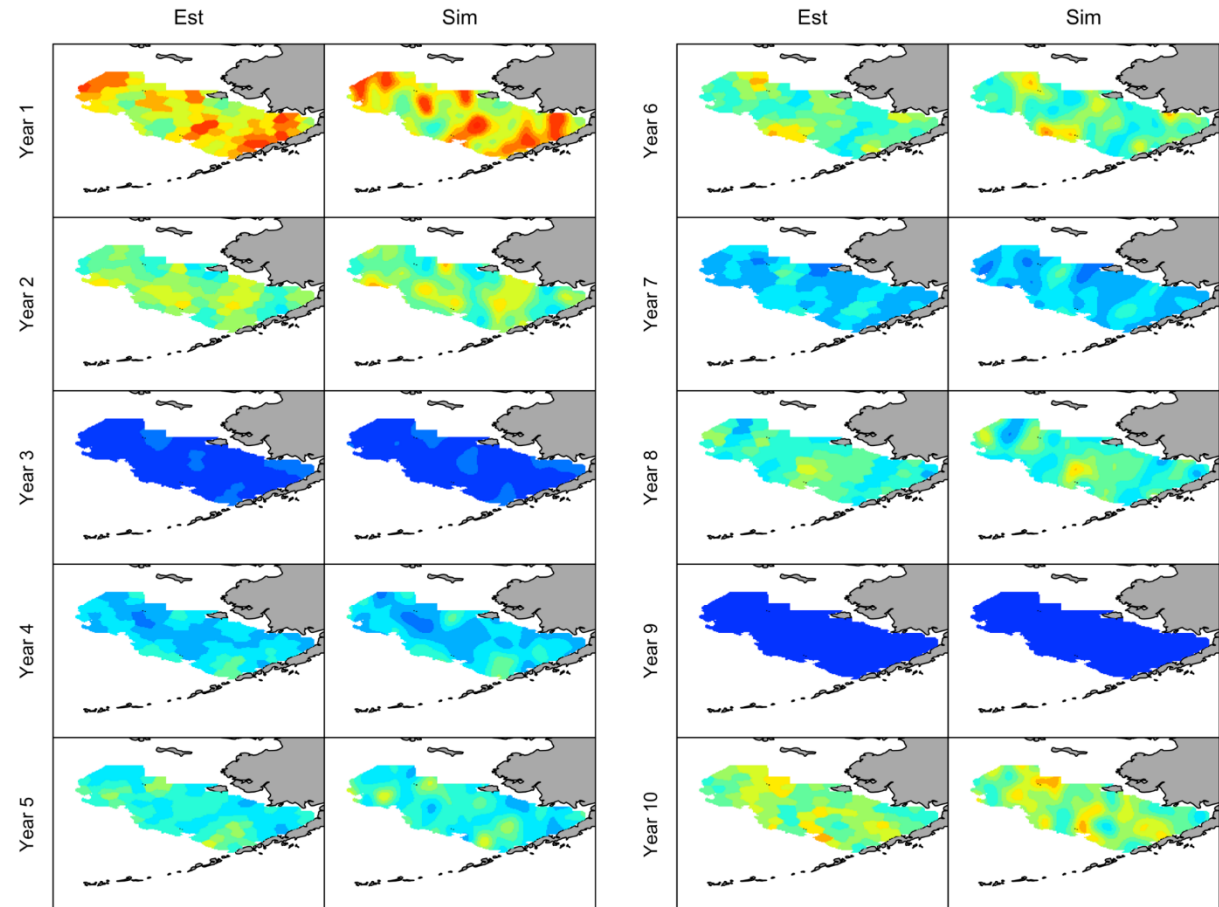
# Simulation experiments

— Experiment 1: Exploring movement

## Catch



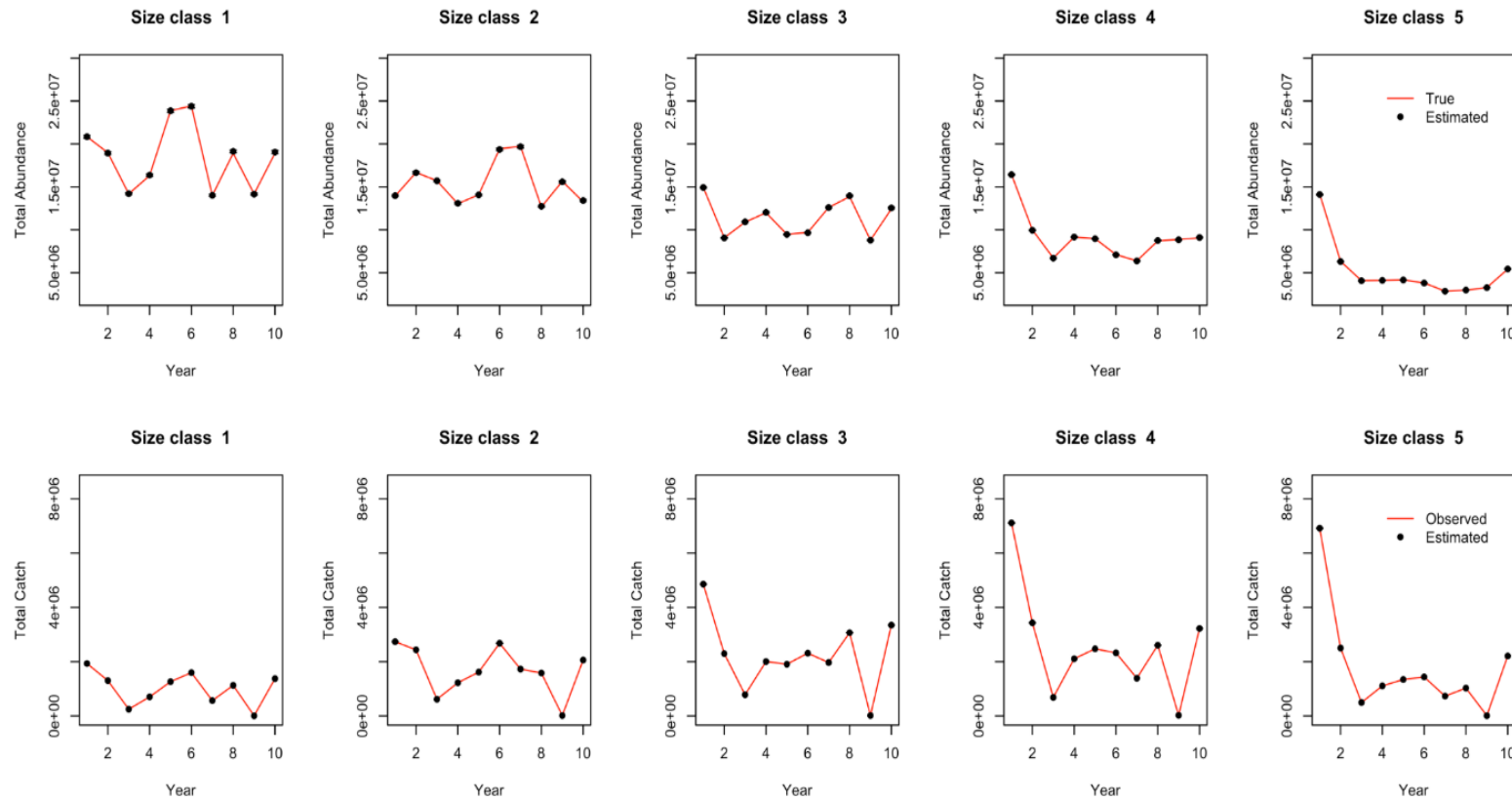
## Fishing mortality



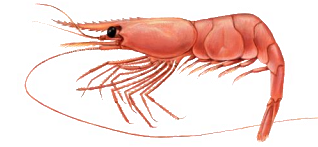
# Simulation experiments

## — Experiment 1: Exploring movement

Spatially-aggregated total abundance (a) and total removals (b) by size class over time

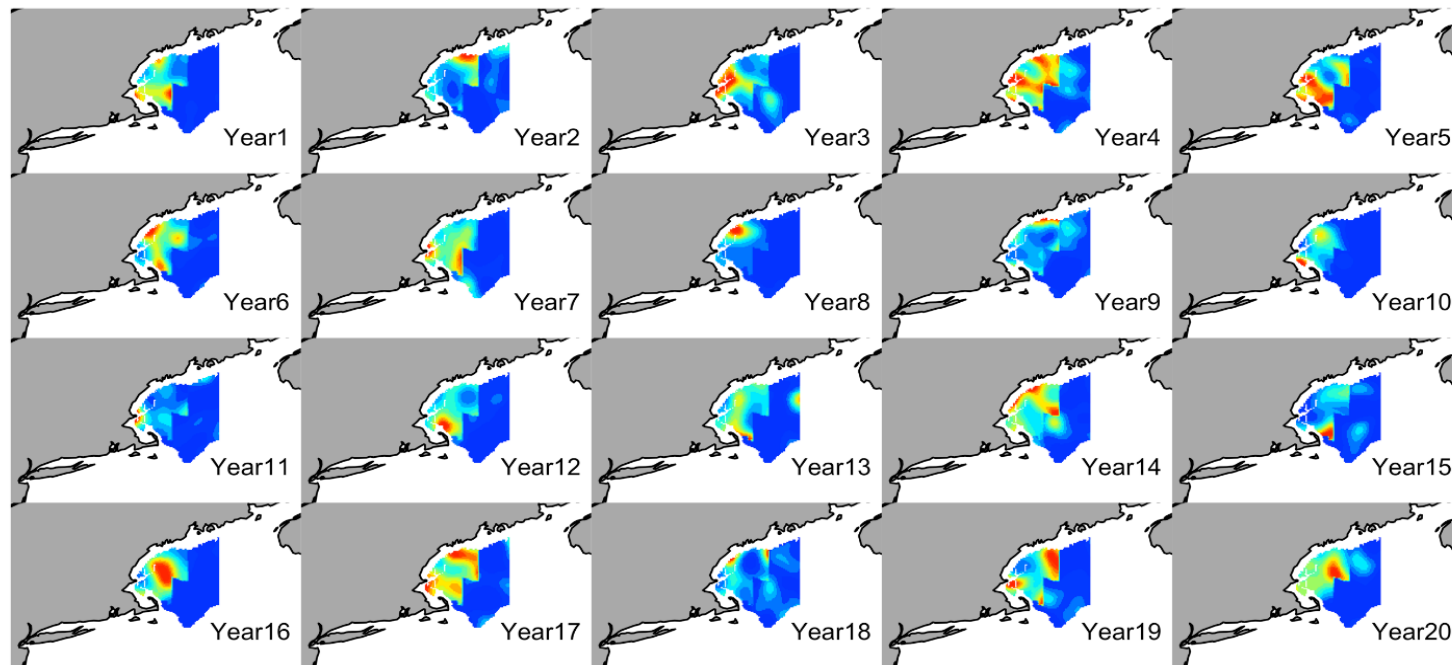


# Simulation experiments



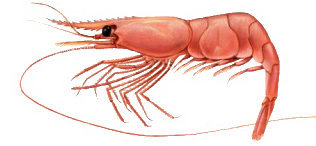
– Experiment 2: comparison of spatiotemporal and spatially-implicit models

Spatially-implicit model – size structured assessment model for *Pandalus* (Cao et al. 2017 )



Simulated fishing mortality for northern shrimp (inshore area has persistent higher fishing mortality over time than offshore area)

# Simulation experiments



– Experiment 2: comparison of spatiotemporal and spatially-implicit models

- the data used in both estimation models are the same at the grid spatial scale
- 50 knots for the spatiotemporal model
- a metric that is directly comparable
- abundance-at-size, fishing mortality at size and spawning stock biomass aggregated over the spatial domain

- **population-level fishing mortality**
- **aggregate selectivity-at-length**

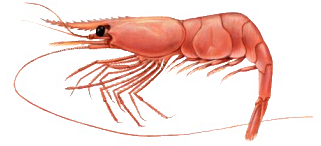
$$c_{l,t} = \left(1 - \exp(-s_{l,t}f_t)\right) n_{l,t} \exp(-m_t)$$

$$\text{RMSE}_l = \sqrt{\frac{\sum_t \left( n_{l,t}^{\text{est}} - n_{l,t}^{\text{true}} / n_{l,t}^{\text{true}} \right)^2}{\tau}} \times 100\%$$

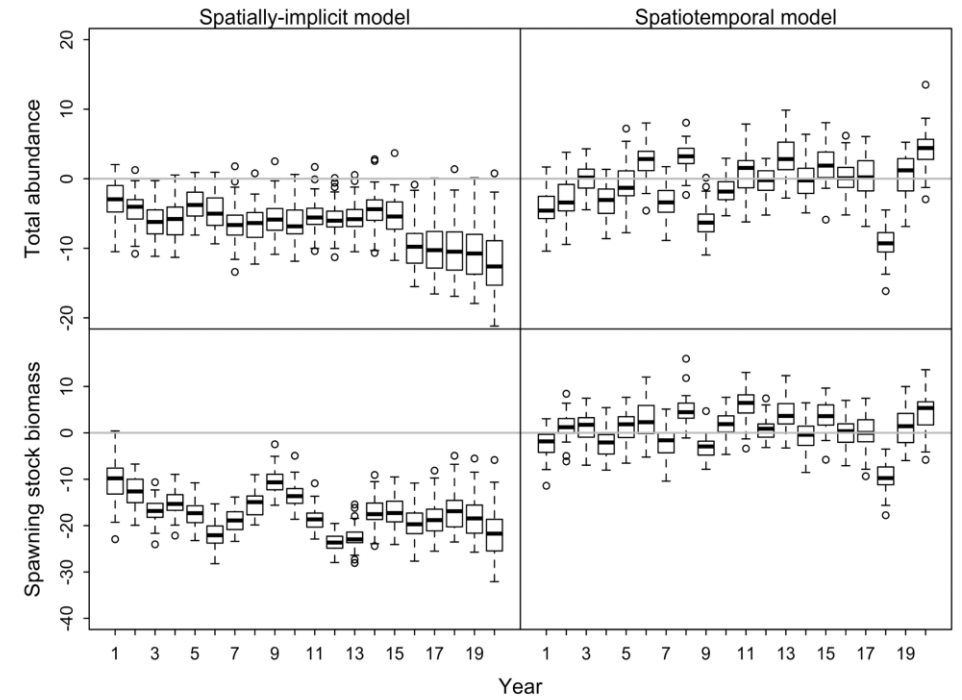
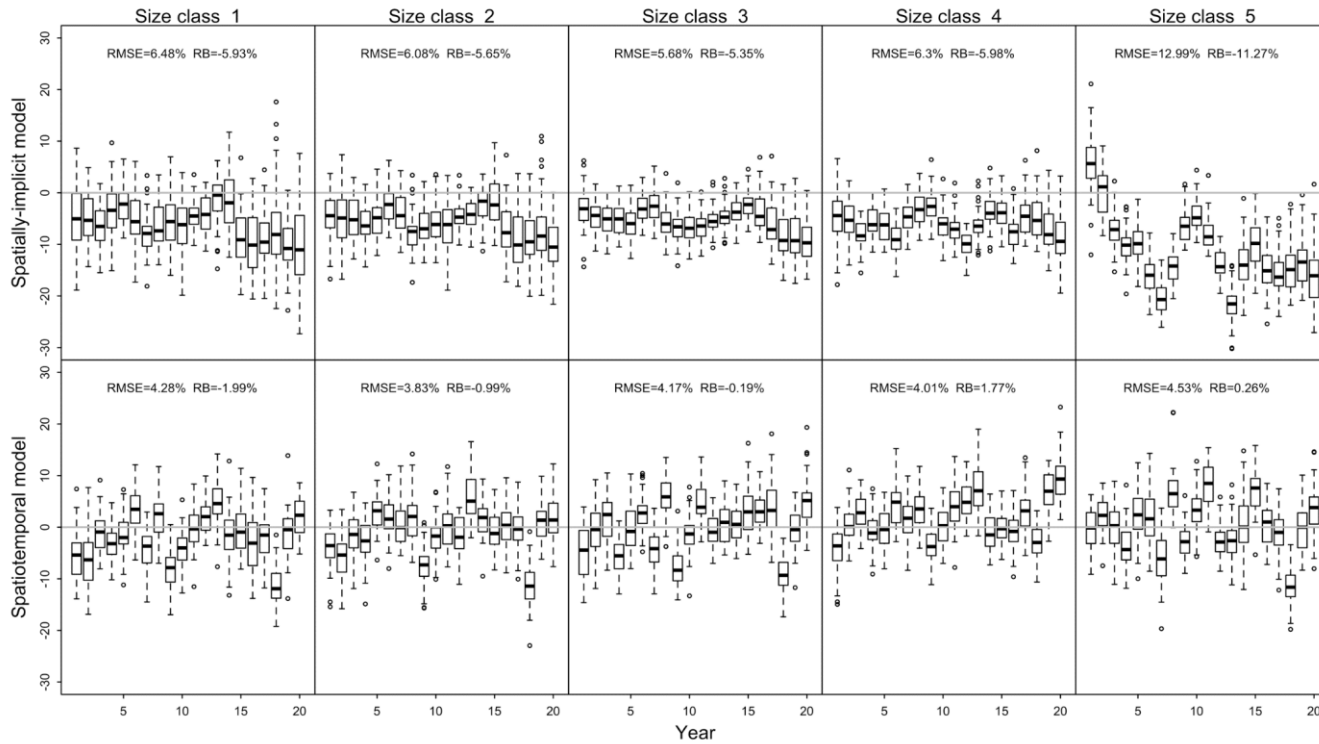
$$\text{RB}_l = \frac{\sum_t \left( n_{l,t}^{\text{est}} - n_{l,t}^{\text{true}} / n_{l,t}^{\text{true}} \right)}{\tau} \times 100\%$$



# Simulation experiments

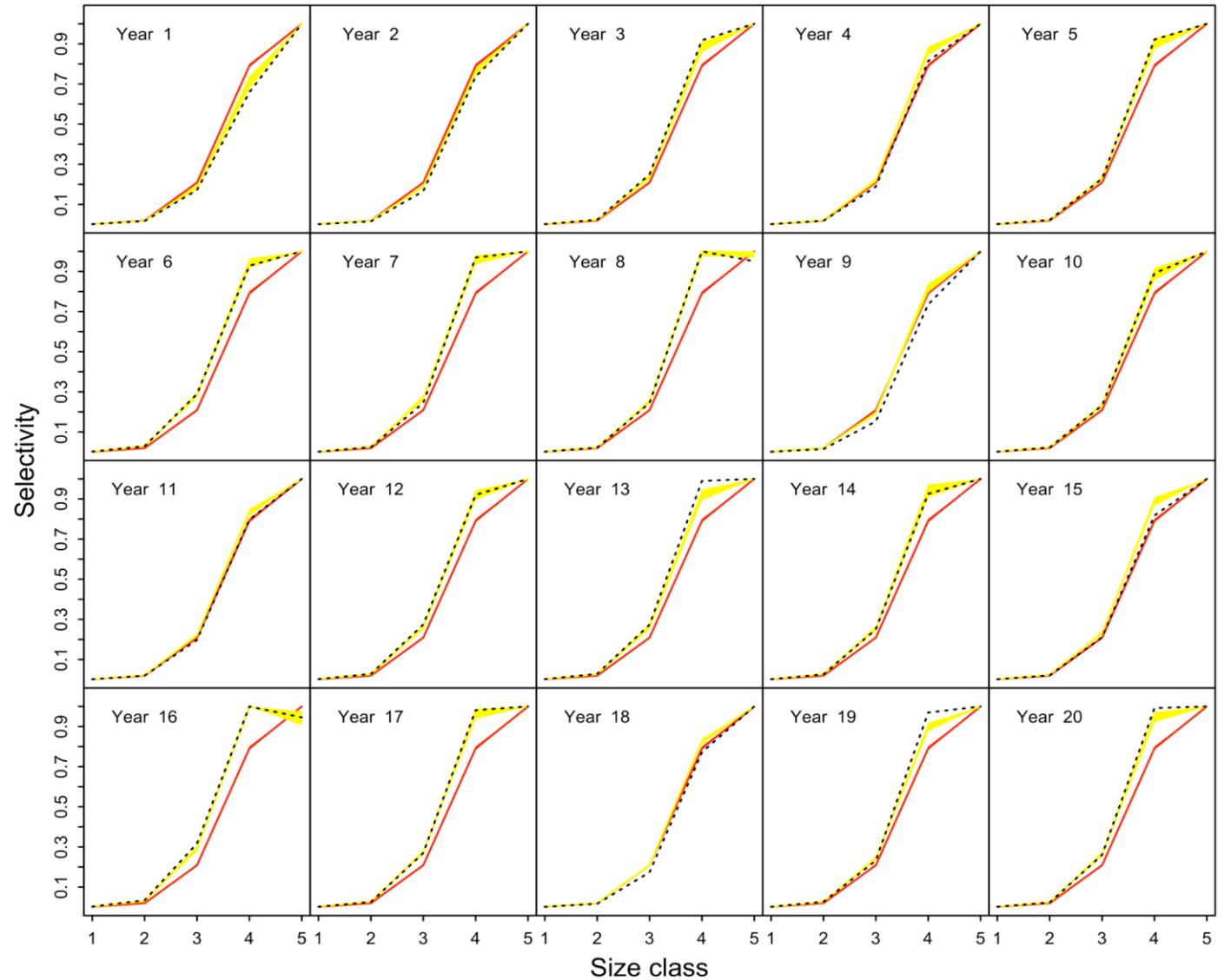


– Experiment 2: comparison of spatiotemporal and spatially-implicit models



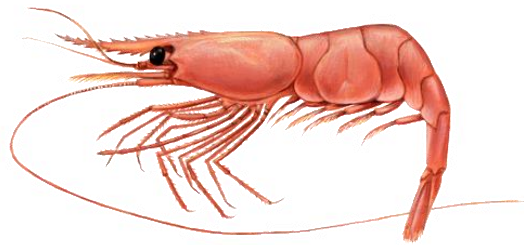
# Simulation experiments

– Experiment 2: comparison of spatiotemporal and spatially-implicit models



# Simulation experiments

— Experiment 3: Effect of sample size



data poor

50 locations

moderate level

100 locations

data rich

200 locations



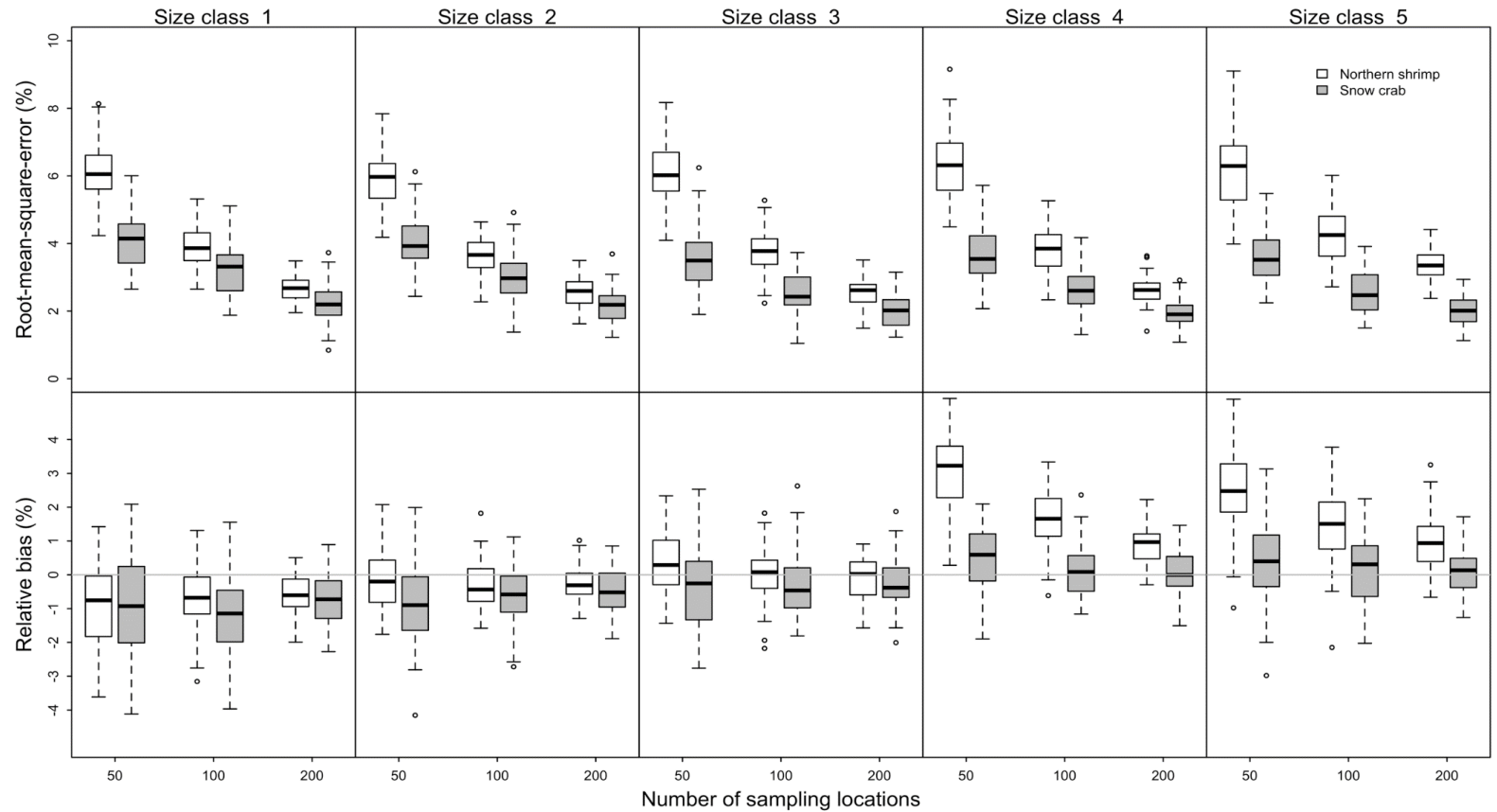
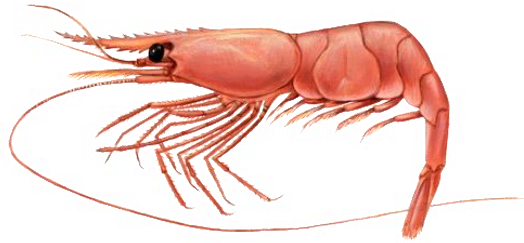
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$$\text{RB}_l = \frac{\sum_t \left( n_{l,t}^{\text{est}} - n_{l,t}^{\text{true}} / n_{l,t}^{\text{true}} \right)}{\tau} \times 100\%$$



# Simulation experiments

— Experiment 3: Effect of sample size



# Conclusions

- The spatiotemporal model produced unbiased estimates of abundance and fishing mortality spatially
- The spatiotemporal model outperformed a spatially-implicit model when time-varying selectivity caused by spatial heterogeneity in fishing pressure is ignored
- Our modeling approach bridges the gap between species distribution and population dynamic models and provides the opportunity to improve natural resource management and conservation

# Discussion

- Adapt to populations with different types of life history through straightforward modifications
- The comparison scenario we show here represents the situation where a strong and persistent gradient of fishing pressure occurs over space and time
- Possible to explicitly model movement
- Selectivity —
  - more biologically interpretable
  - could be corroborated by other field sampling
- The advance comes at the expense of greater data requirements

# ACKNOWLEDGEMENTS

- We thank K. Kristensen and the developers of Template Model Builder, without which this analysis would not be feasible
- NOAA “Stock Assessment and Analytic Methods” (SAAM) grant
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